

```
data ex2018_2_10_1;
```

```
input species $ gender $ amino@@; output;
```

```
datalines;
```

```
m1 m 21.5 m1 m 19.6 m1 m 20.9 m1 m 22.8 m1 f 14.8 m1 f 15.6 m1 f  
13.5 m1 f 16.4
```

```
m2 m 14.5 m2 m 17.4 m2 m 15.0 m2 m 17.8 m2 f 12.1 m2 f 11.4 m2 f  
12.7 m2 f 14.5
```

```
m3 m 16.0 m3 m 20.3 m3 m 18.5 m3 m 19.3 m3 f 14.4 m3 f 14.7 m3 f  
13.8 m3 f 12.0
```

```
;
```

```
proc glm;
```

```
class species;
```

```
model amino=species;
```

```
means species/ hovtest=bartlett welch;
```

```
run;
```

```
ods graphics on;
```

```
proc glm data=ex2018_2_10_1 plot=diagnostics;
```

```
class species;
```

```
model amino = species;
```

```
run;
```

```
ods graphics off;
```

```
proc glm;
```

```
class species gender;
```

```
model amino=gender species;
```

```
means species/snk scheffe;
```

```
run;
```

```
ods graphics on;
```

```
proc glm data=ex2018_2_10_1 plot=diagnostics;
```

```
class species gender;
```

```
model amino = species gender;
```

```
run;
```

```
ods graphics off;
```

```

data ex2018_2_10;
do species='m1', 'm2', 'm3';
do gender='male', 'female';
do k=1 to 4;
    input amino@@; output;
    end; end; end;
datalines;
21.5 19.6 20.9 22.8 14.8 15.6 13.5 16.4
14.5 17.4 15.0 17.8 12.1 11.4 12.7 14.5
16.0 20.3 18.5 19.3 14.4 14.7 13.8 12.0
;
proc glm;
class species gender;
model amino=species gender species*gender;
run;
proc glm;
class species gender;
model amino=species gender;
run;
proc glm;
class species gender;
model amino=gender species;
means species/ tukey duncan;
run;

```

```

data ex3_1;
  input drug $ result $ count;
  datalines;
  aspirin my_y 139
  aspirin my_n 10898
  placebo my_y 239
  placebo my_n 10795
  ;
proc freq;
  tables drug*result/chisq measures;
  weight count;
run;

```

```

data ex3_2;
  input drug $ result $ count;
  datalines;
  yes my_y 73
  yes my_n 18
  no my_y 141
  no my_n 196
  ;
proc freq;
  tables drug*result/chisq measures;
  weight count;
run;

```

```

data ex3_2_1;
  input drug $ result $ count;
  datalines;
  drug m_d 73
  drug m_n 18
  no m_d 141
  no m_n 196
  ;
proc freq;
  tables drug*result/chisq measures;
  weight count;
run;

```

```

data ex3_5;
  input before $ after $ count;
  datalines;
sat sat 23
sat uns 7
uns sat 18
uns uns 12
;
proc freq;
  tables before*after/chisq agree;
  weight count;
run;

```

```

DATA ex3_6;
INPUT hospital $ trt $ recovery $ count @@;
CARDS;
A old yes 9 A old no 5
A new yes 11 A new no 6
B old yes 7 B old no 5
B new yes 8 B new no 3
C old yes 4 C old no 6
C new yes 7 C new no 5
D old yes 18 D old no 11
D new yes 26 D new no 4
;
PROC FREQ;
WEIGHT count;
TABLES trt*recovery/chisq;
RUN;
PROC FREQ;
WEIGHT count;
TABLES hospital*trt*recovery/CMH NOROW NOCOL;
RUN;
PROC FREQ;
WEIGHT count;
TABLES hospital*trt*recovery/CMH1 NOROW NOCOL;
RUN;

```